

#6

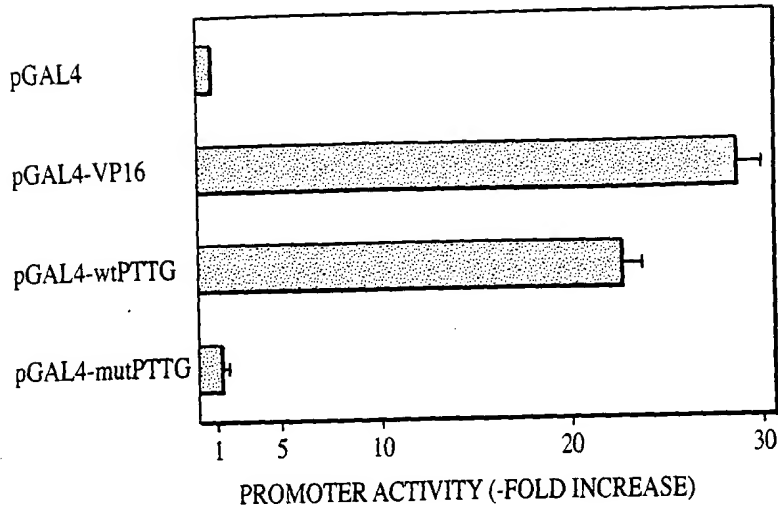


FIG. 1

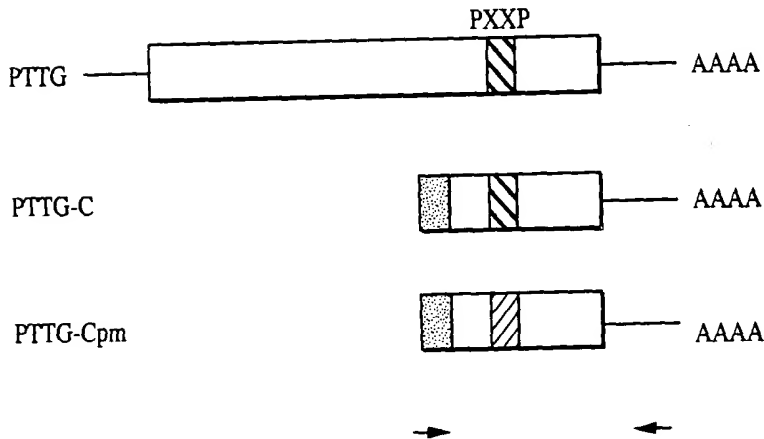


FIG. 2A

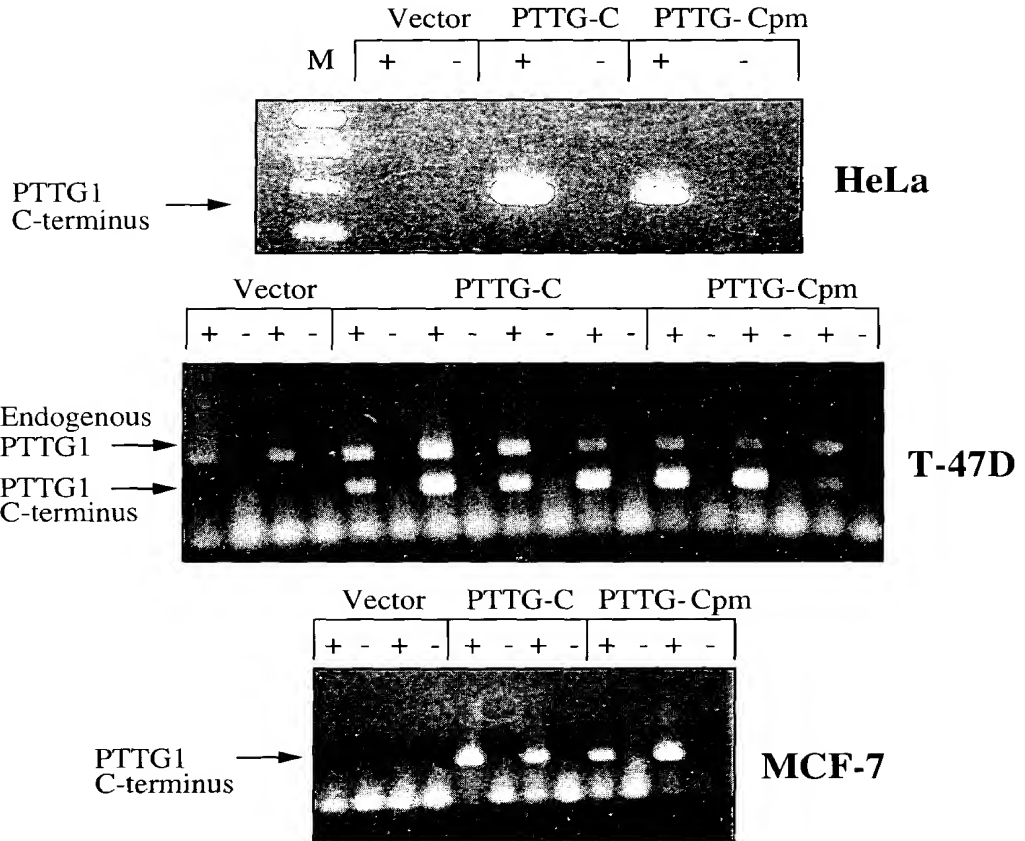


FIG. 2B

001250" 92E15860

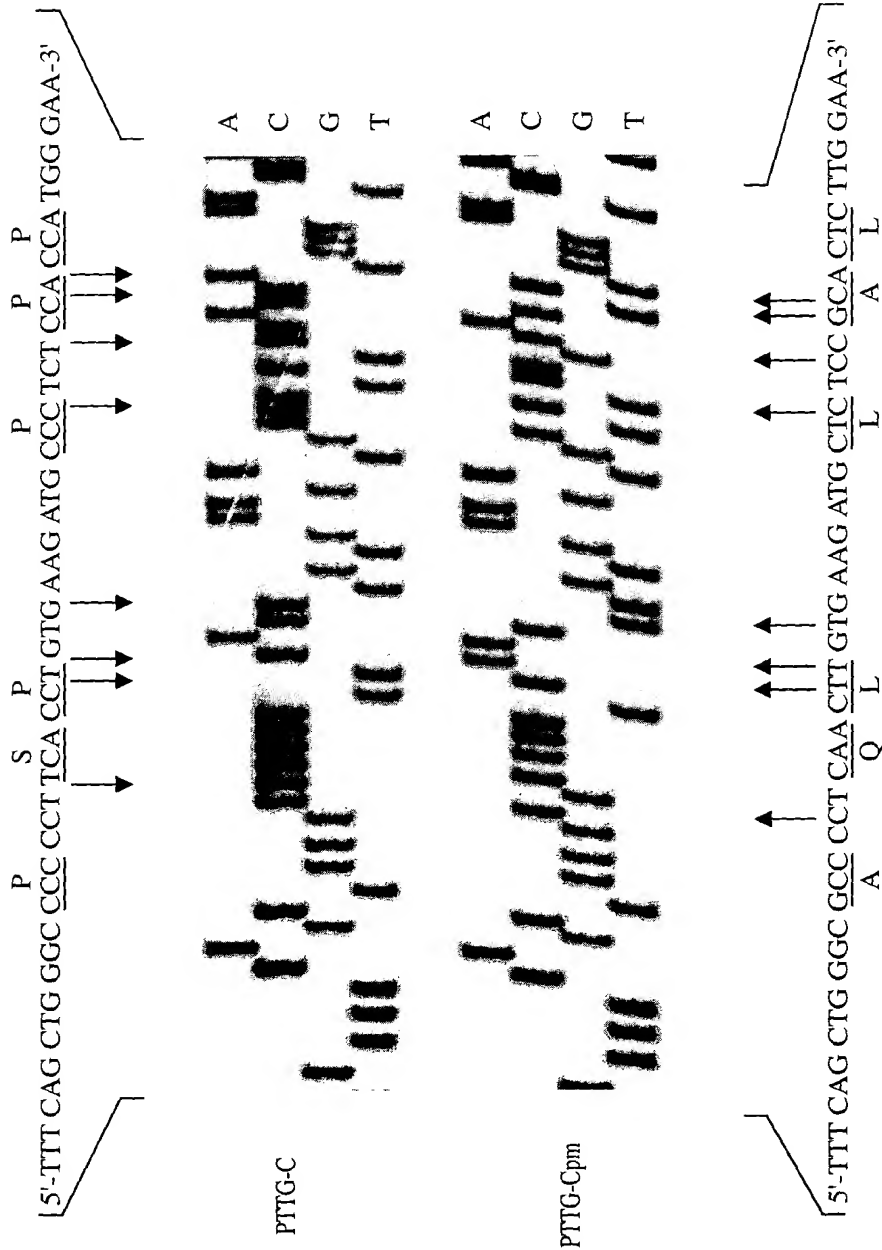
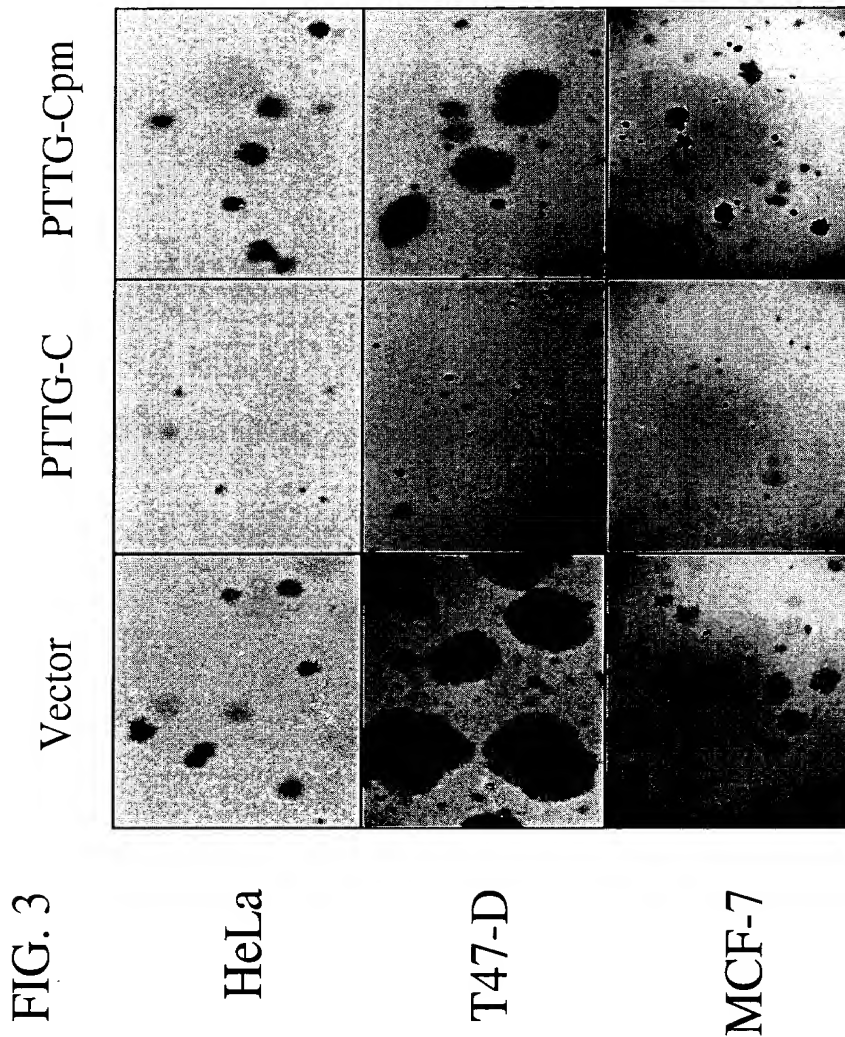


FIG. 2C

107250-9254555



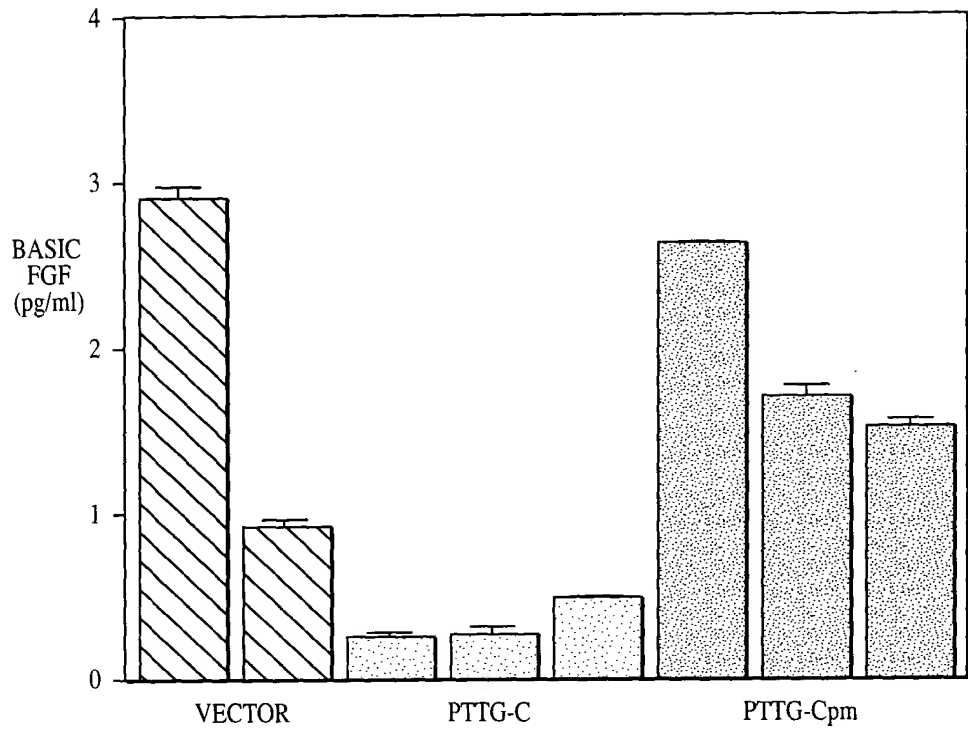


FIG. 4

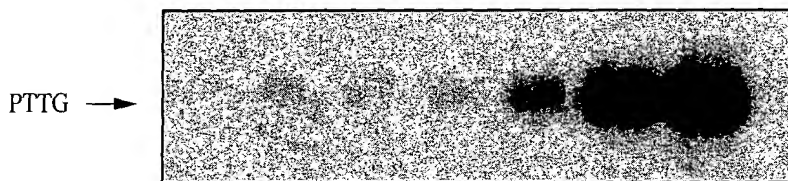
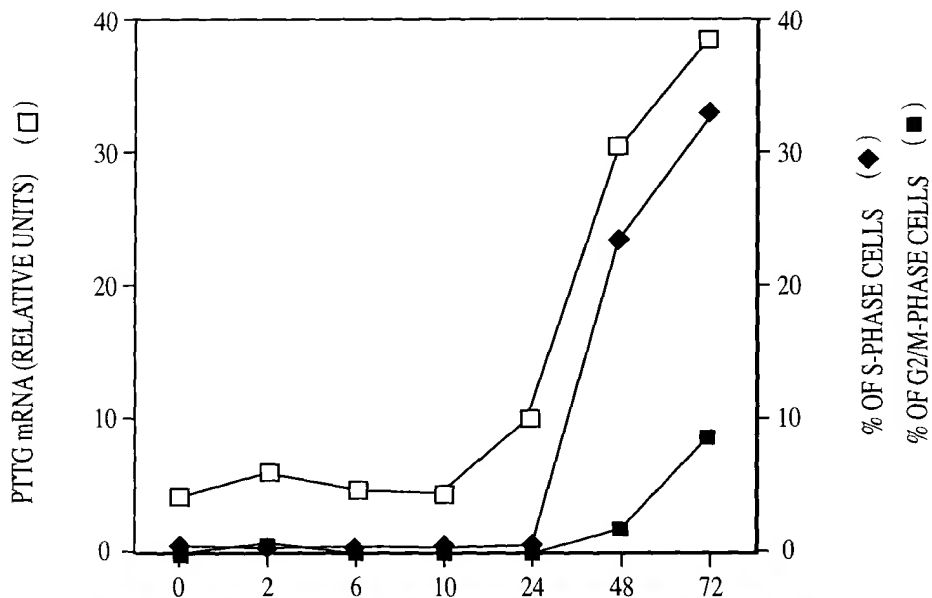


FIG. 5

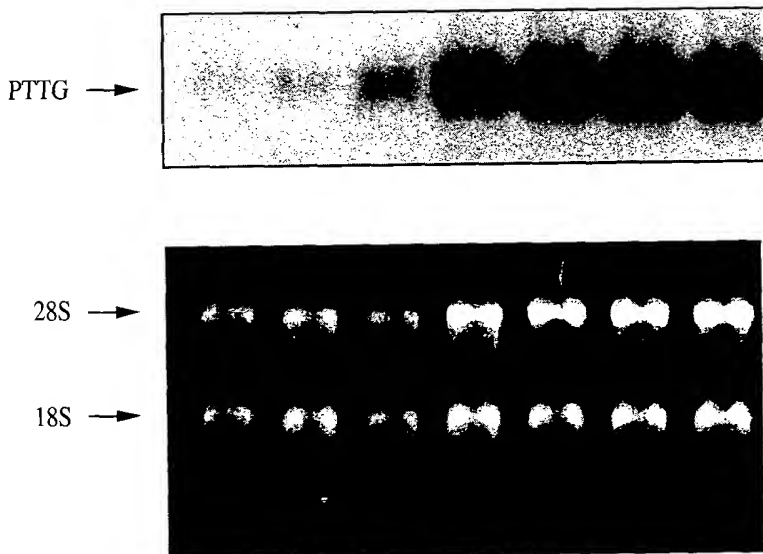
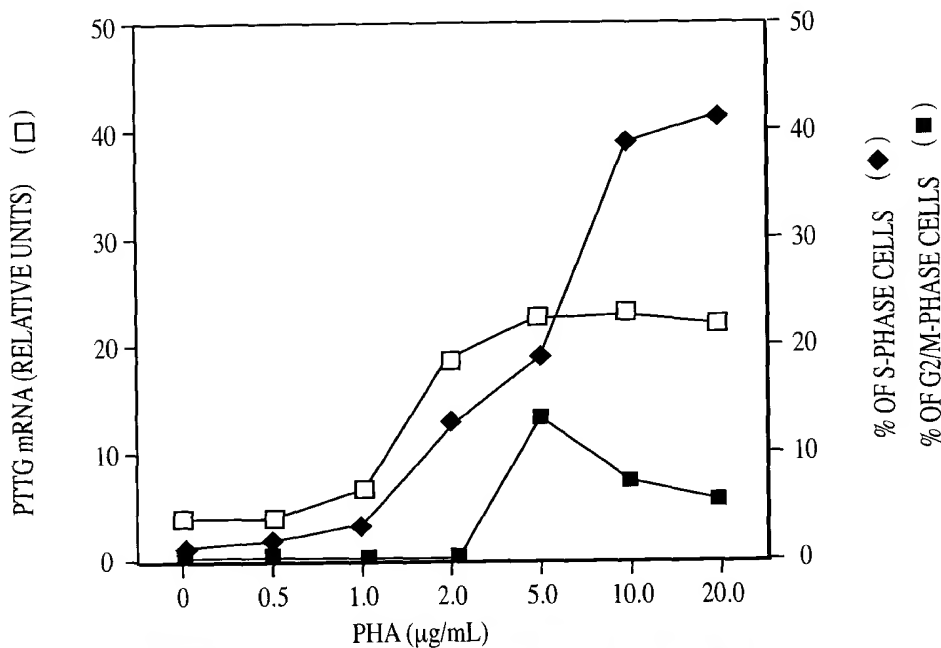


FIG. 6

0054326.002104

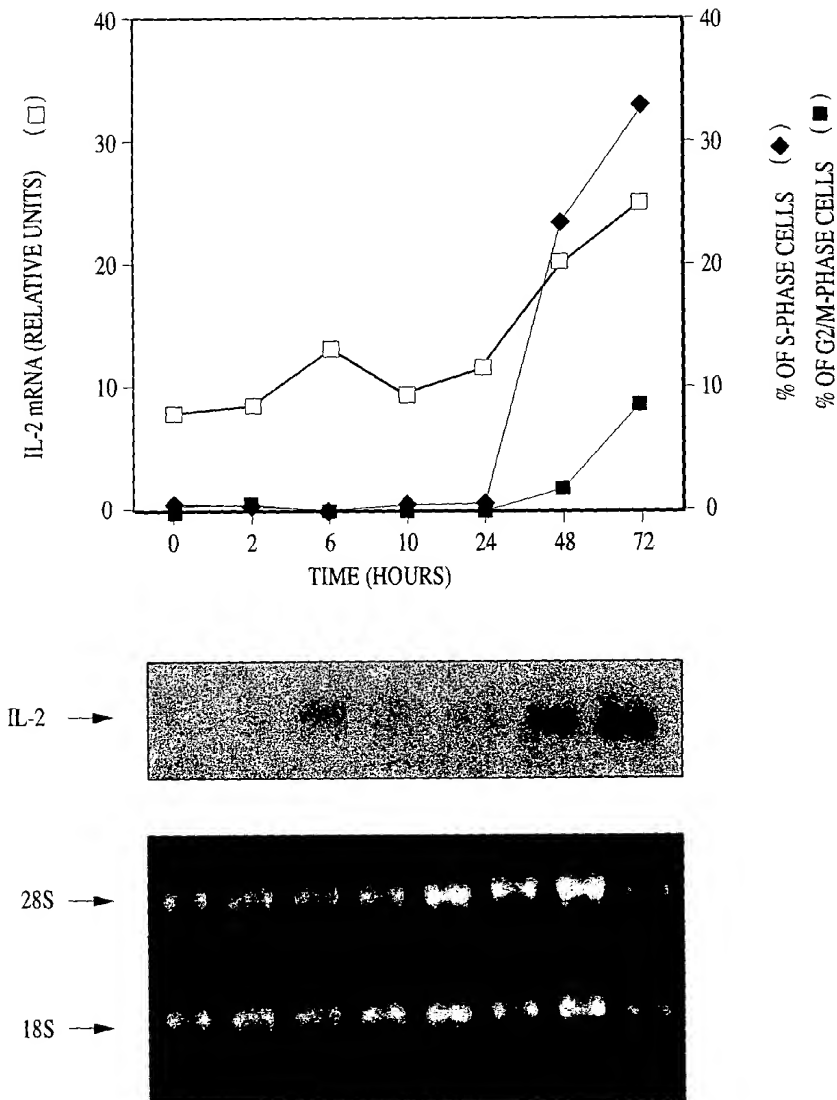


FIG. 7

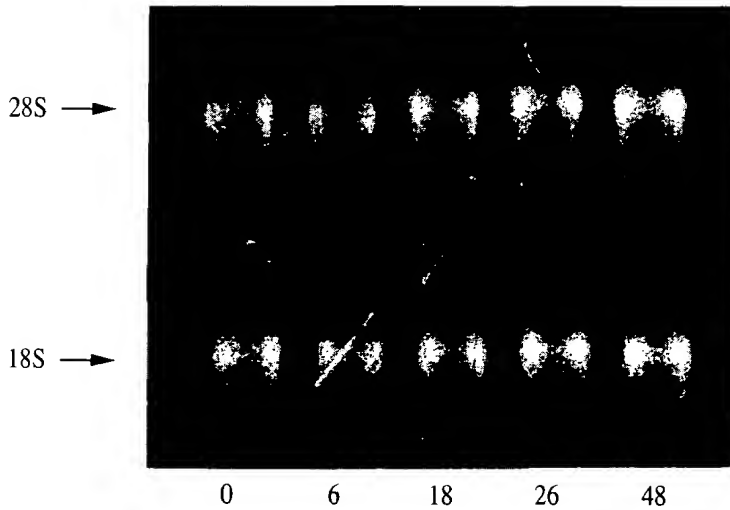
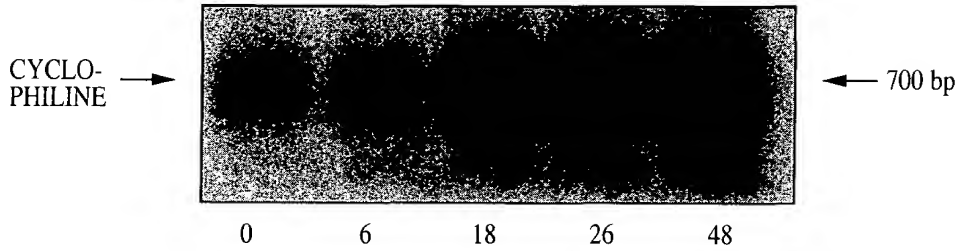
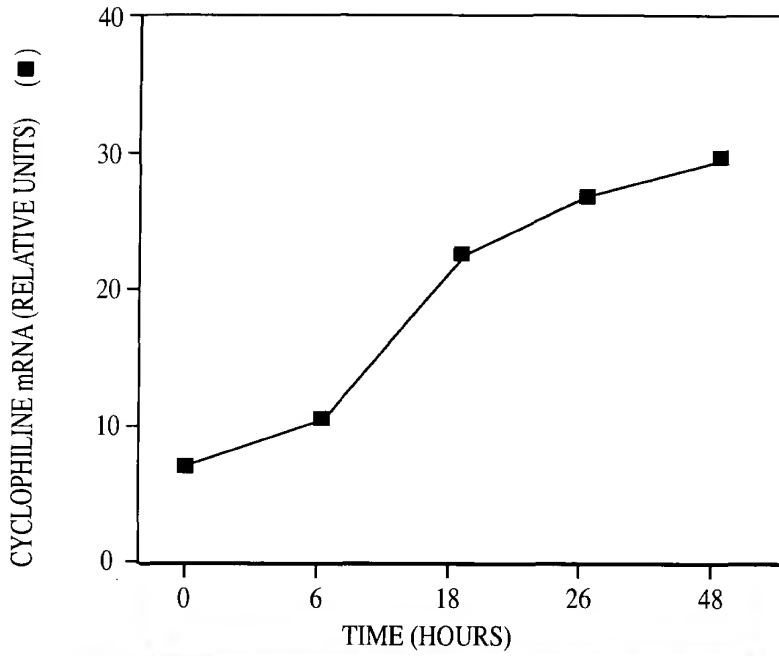


FIG. 8

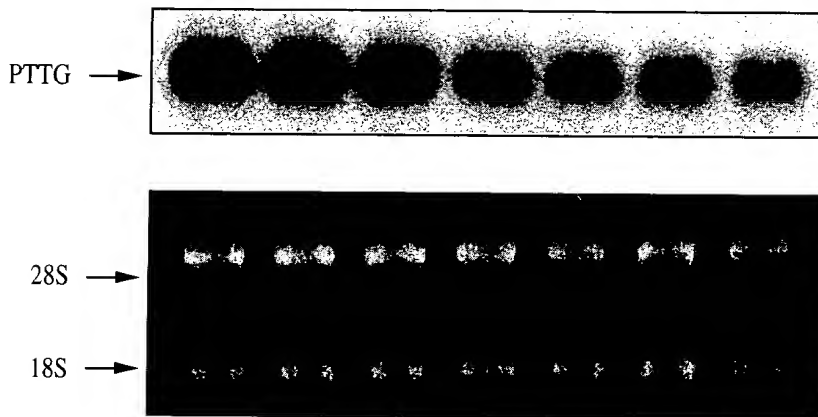
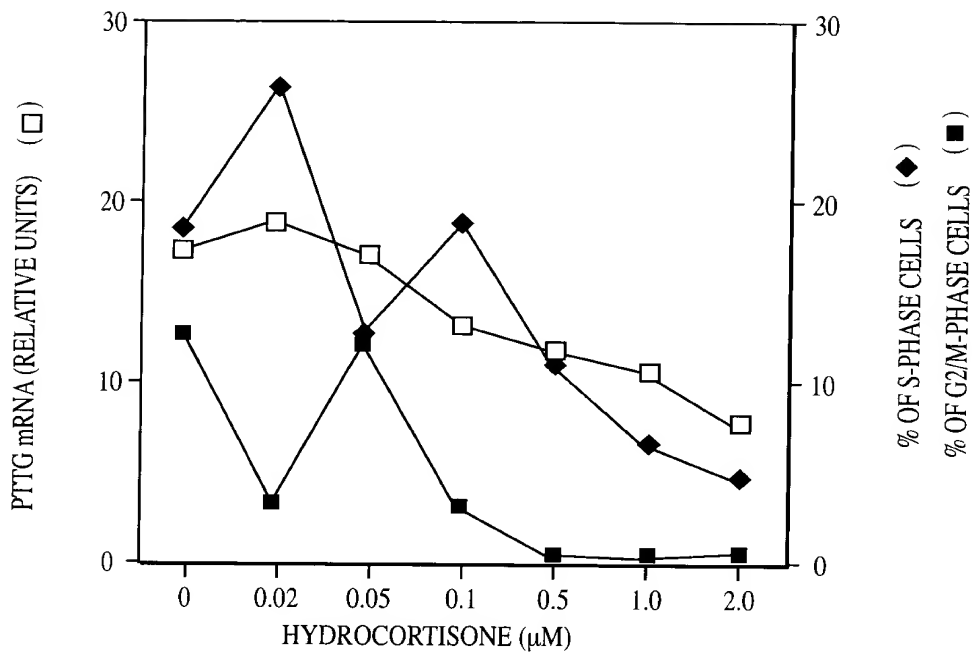


FIG. 9

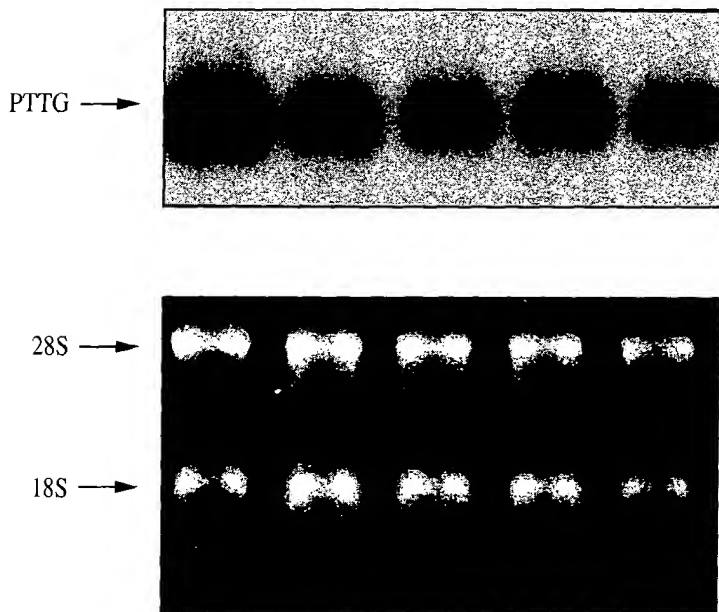
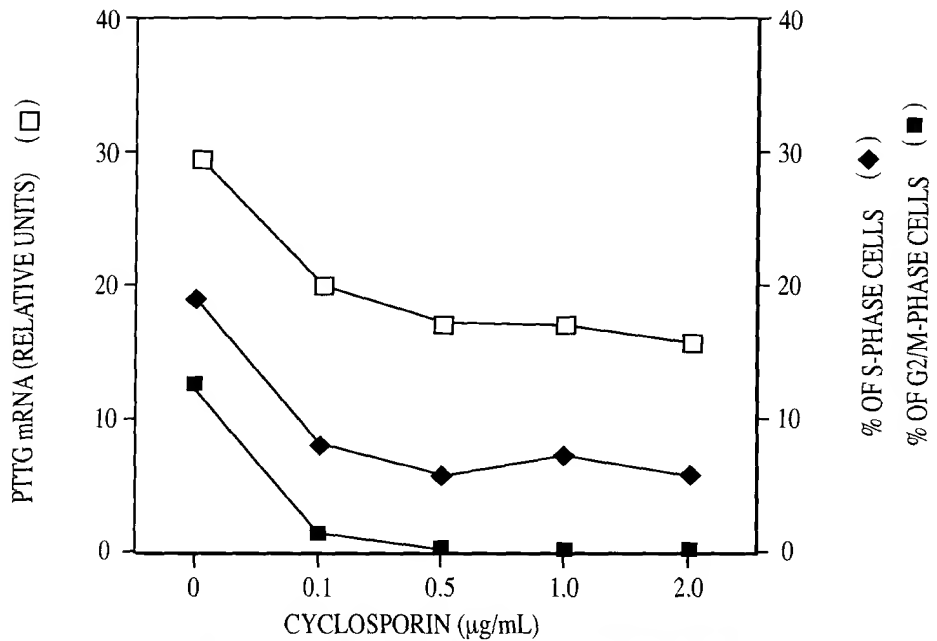


FIG. 10

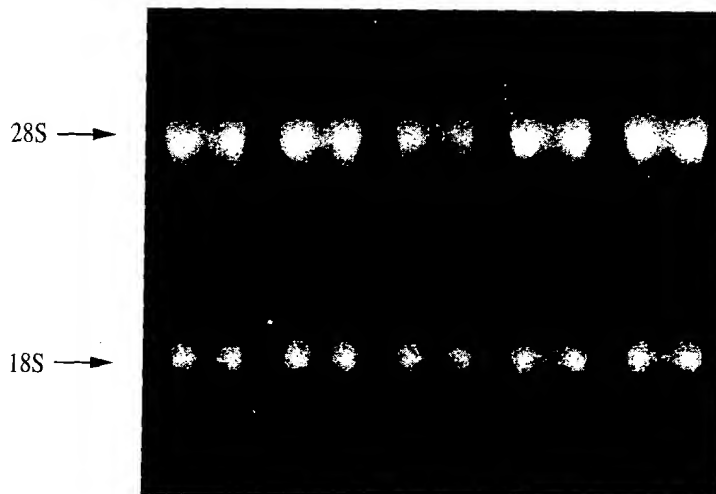
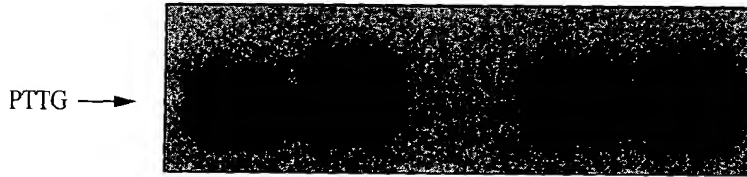
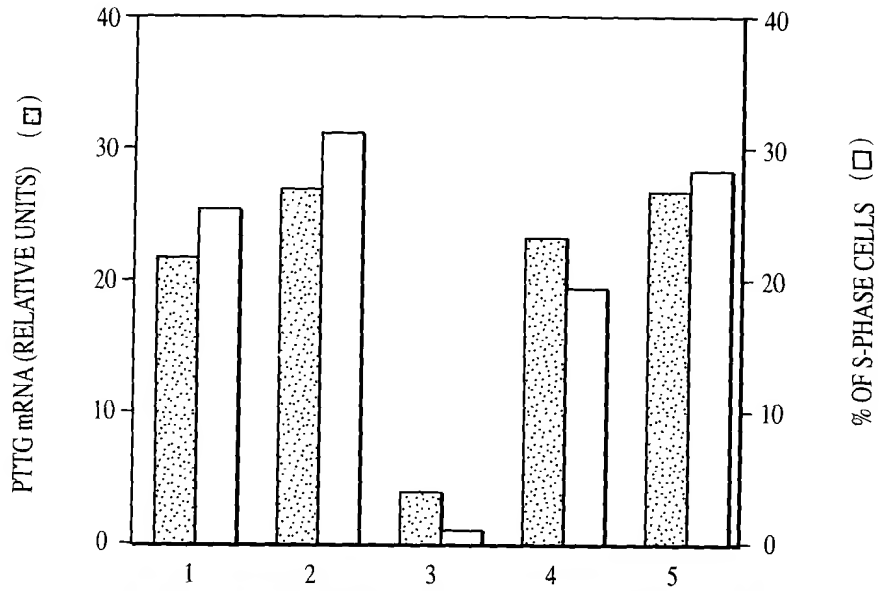


FIG. 11

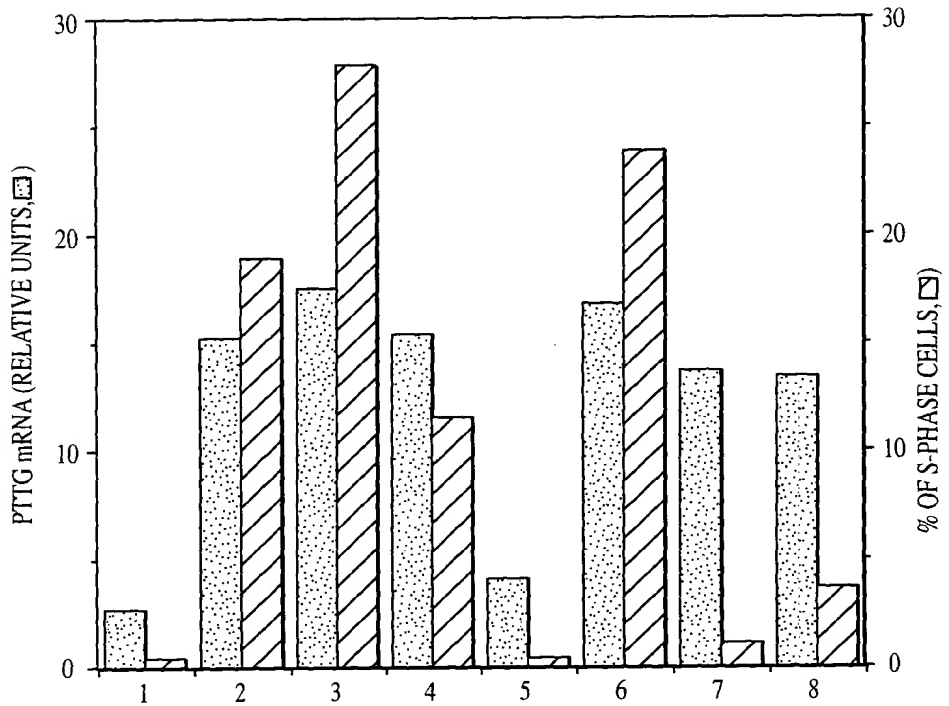


FIG. 12

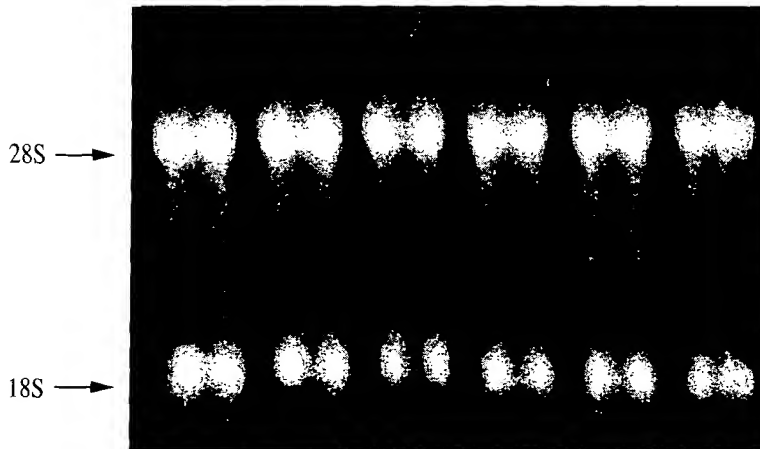
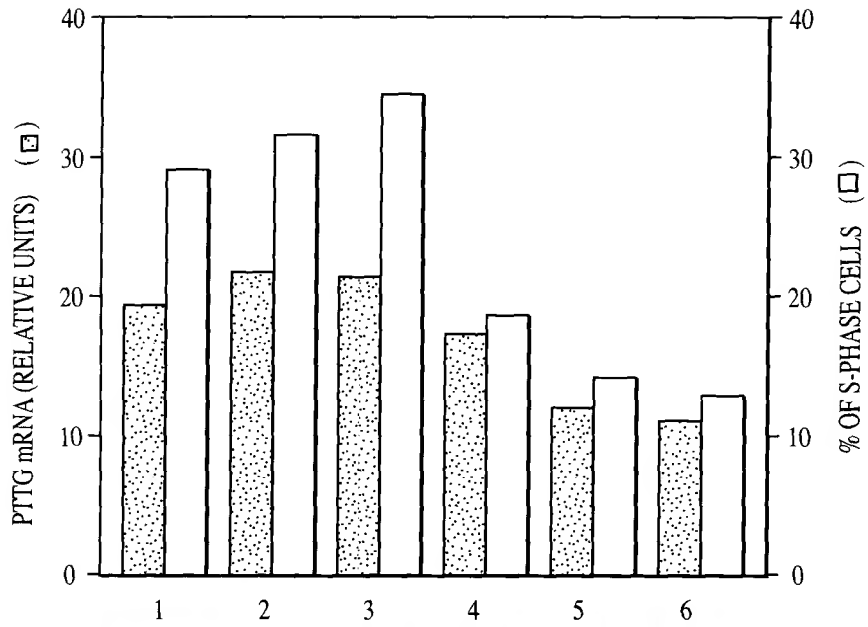


FIG. 13

107260* 22515360

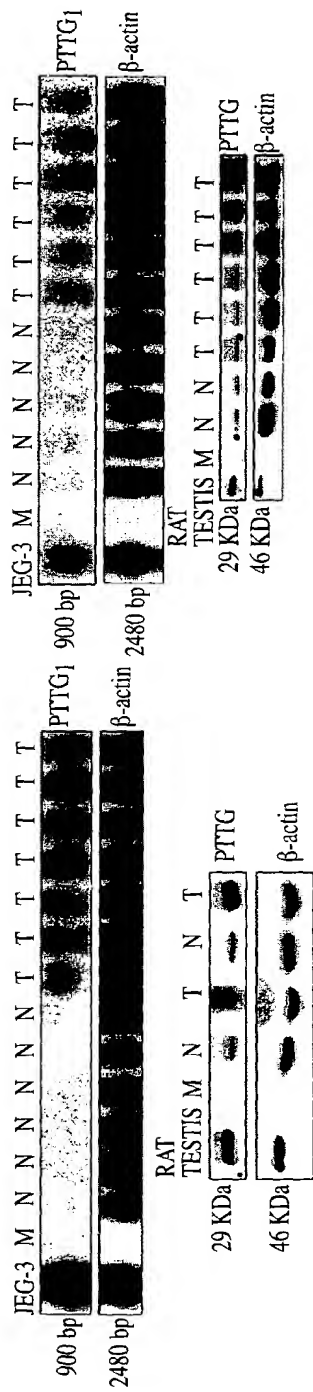


FIG. 14A

FIG. 14B

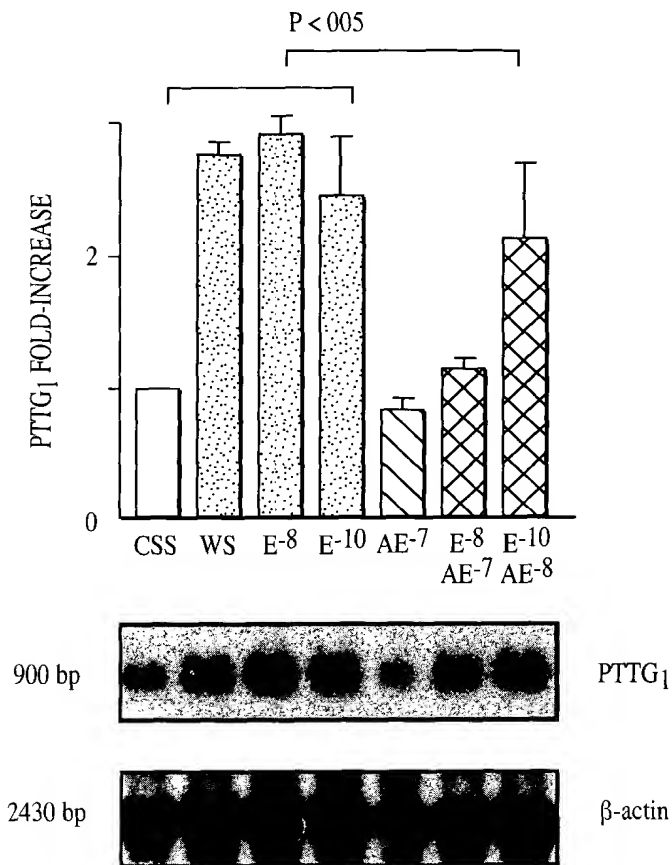


FIG. 15A

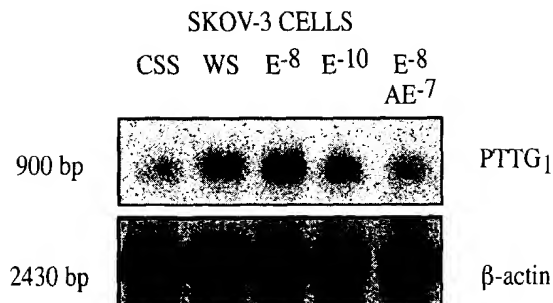


FIG. 15B

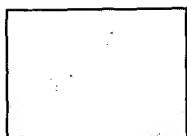


FIG. 16A

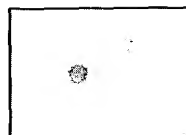


FIG. 16E



FIG. 16B

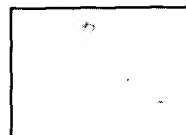


FIG. 16F



FIG. 16C



FIG. 16G



FIG. 16D



FIG. 16H

FIG. 16A, 16B, 16C, 16D, 16E, 16F, 16G, 16H

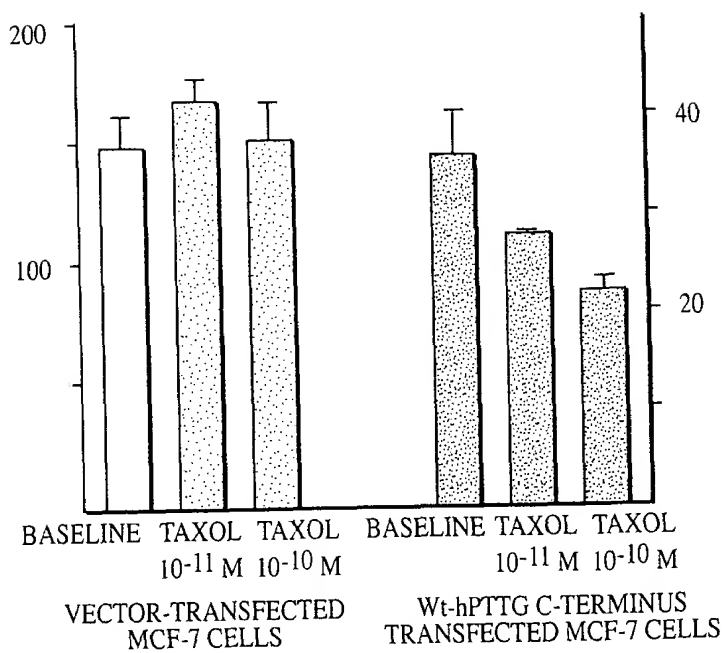


FIG. 17

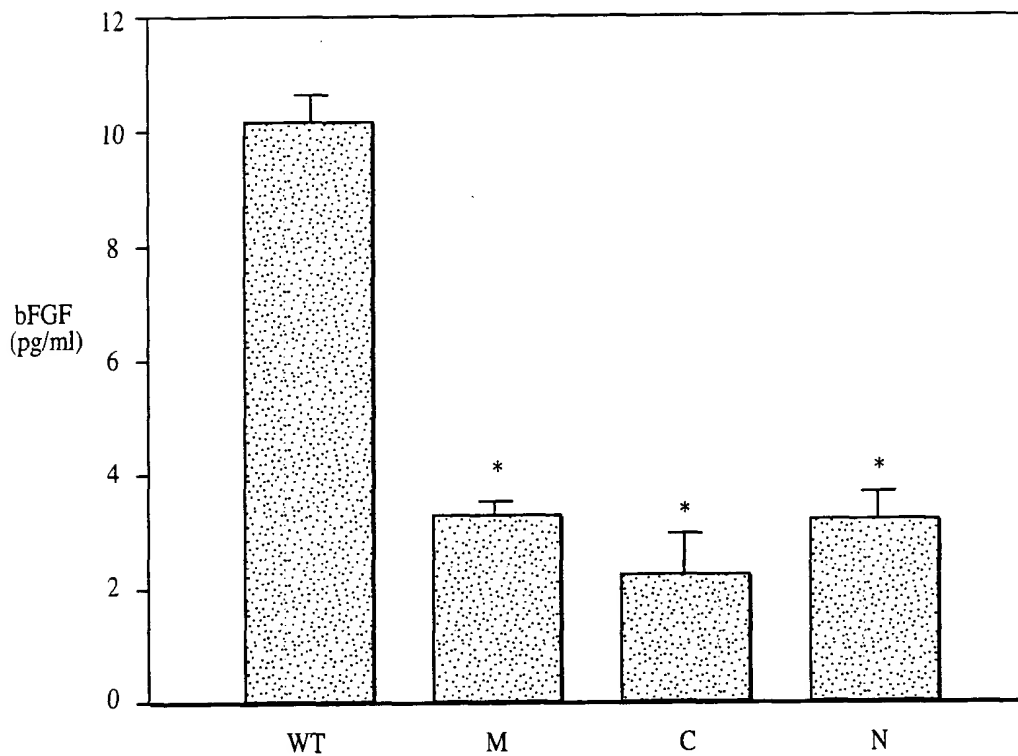


FIG. 18

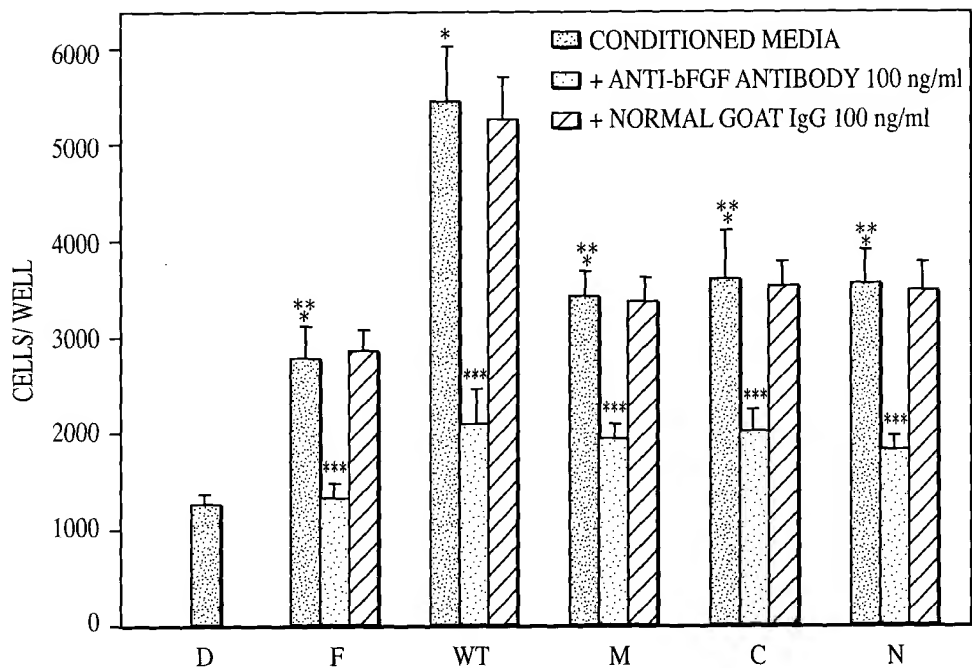


FIG. 19

101250 22545860

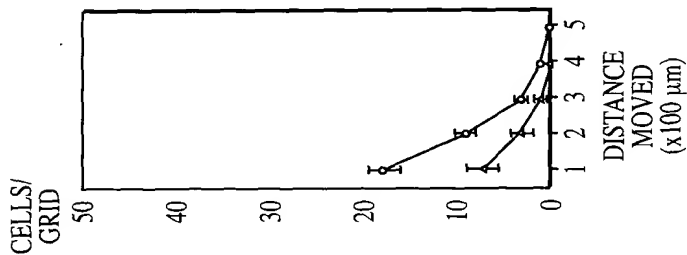


FIG. 20B-a

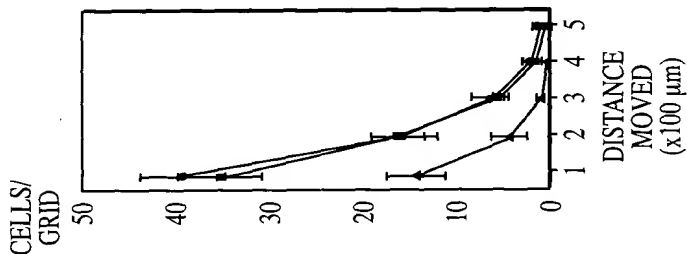


FIG. 20B-b

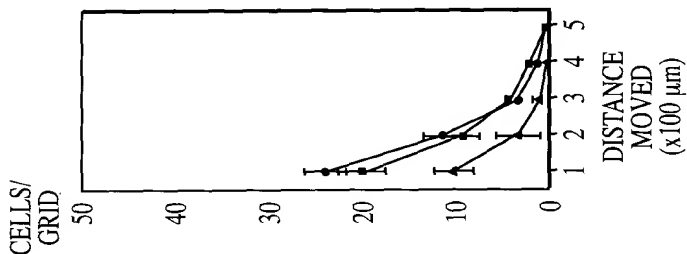


FIG. 20B-c

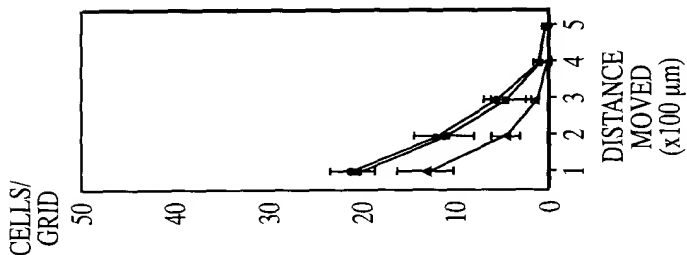


FIG. 20B-d

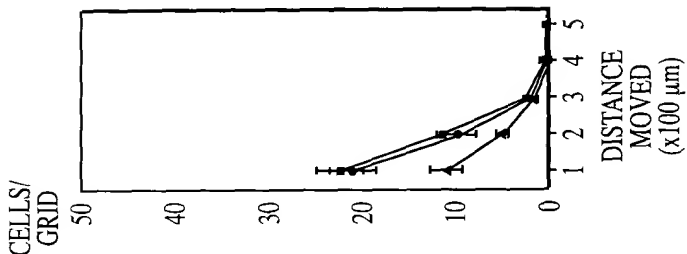


FIG. 20B-e

107260* 92545860

200 μ m



FIG. 21A-a

200 μ m

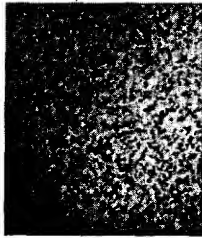


FIG. 21A-b

200 μ m

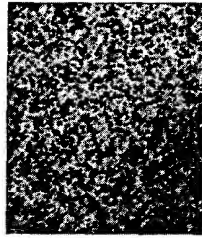


FIG. 21A-c

200 μ m

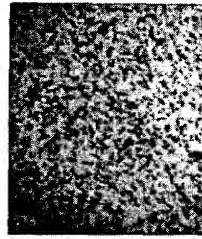


FIG. 21A-d

200 μ m

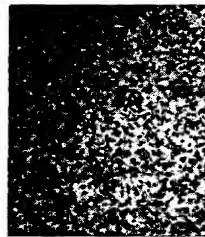


FIG. 21A-e

200 μ m

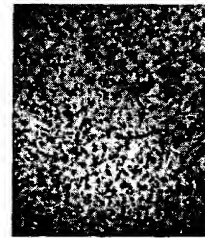


FIG. 21A-f

200 μ m

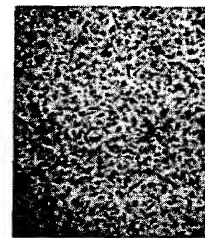


FIG. 21A-g

200 μ m

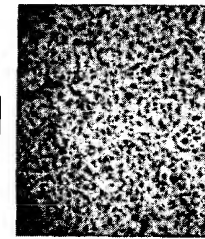


FIG. 21A-h

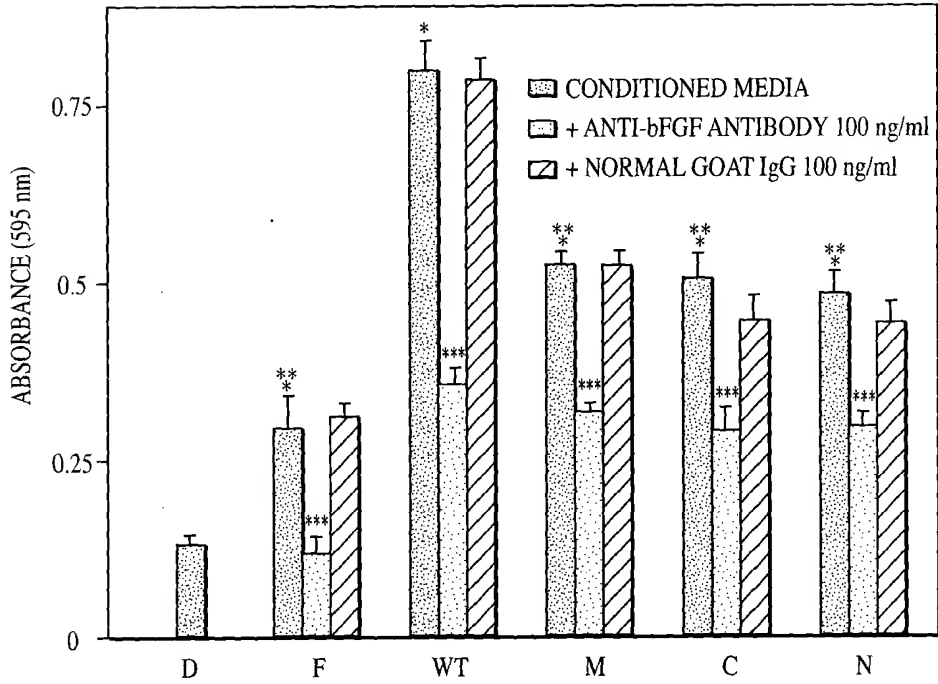


FIG. 21B

101200 9254580

200 μ m



FIG. 22A-a

200 μ m

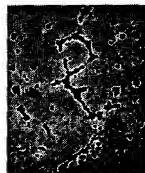


FIG. 22A-b

200 μ m



FIG. 22A-c

200 μ m

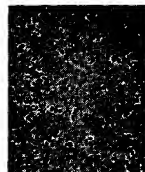


FIG. 22A-d

200 μ m

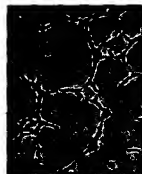


FIG. 22A-e

200 μ m

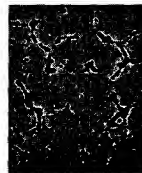


FIG. 22A-f

200 μ m



FIG. 22A-g

200 μ m

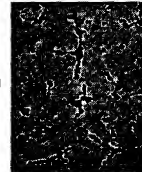


FIG. 22A-h

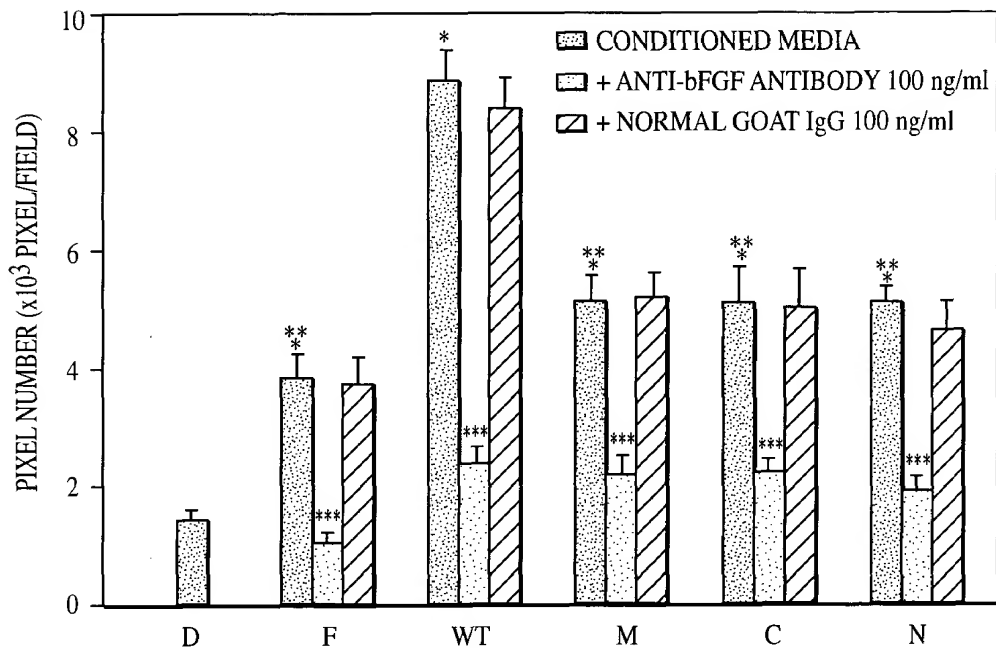


FIG. 22B

1mm



FIG. 23A-a

1mm



FIG. 23A-b

1mm

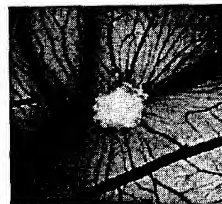


FIG. 23A-c

1mm



FIG. 23A-d

1mm



FIG. 23A-e

1mm

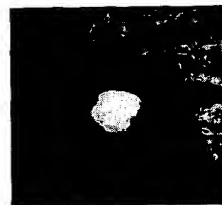


FIG. 23A-f

09/854,326

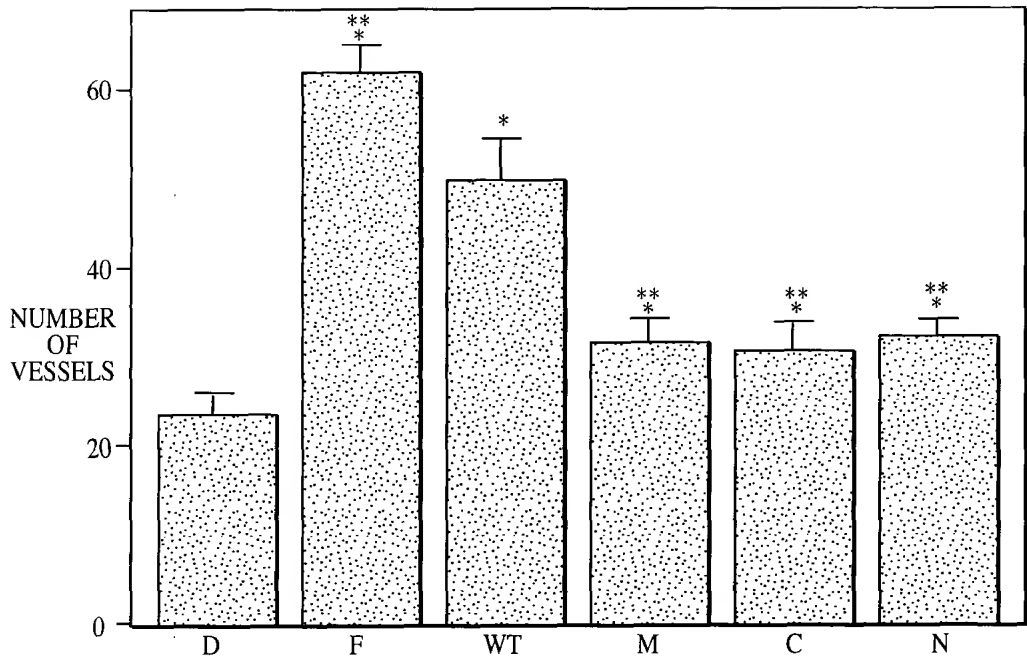


FIG. 23B

hPTTG1	1	MATLIYVDKE	NGEPGTRVVA	KDGLKLGSGP	SIKALDGRSQ	VSTPRFGKTF	DAPPALPKAT
hPTTG2	1	*****	I*****	**V**E**R*	*****I**	*L*****Y	***S*****
hPTTG3	1	*****	*E**IL*AT	*****	*****	**ISC*****	***TS*****
mPTTG	1	*****F**D	*E**R*IAS	*****T*V	--*****KL*	*****V**V*	N*-**V*****S
rPTTG	1	*****F**D	*E**S*IAS	*****V	--*****KL*	*****V**V*	G*-*GL*****S
	61	<u>RKALGTVNRA</u>	TEKSVKTKGP	LKQKQPSFSA	KKMTEKTVKA	KSSVLASDDA	YPEIEKFFPF
	61	*****	*****R	*****	*****T	*****P*****	*****L****
	61	*****	*****	*****	*****	*N**P*****G	*****L****
	58	*****V	A**PM**GK*	*QP***TLTG	**I***ST*T	Q***P**P***	*****L****
	58	*****V	***P**SSK*	*QS***TL*V	**I***ST*T	QG*AP*P***	*****L****
	121	<u>NPLDFESFDL</u>	PEEHQIAHLP	LSGVPLMILD	EERELEKLFQ	<u>LGPPSPVKMP</u>	<u>SPPWESNLLQ</u>
	121	*L*****	*****	*****	**G*****	*****C**FA	*****K*****
	121	*G*****	*****	**E*****	*****L***	*****L***	*****K*****
	118	***--***	*****SL**	*N***IT*N	***G*****LH	*****L*T*	FLS*****DP*Y
	118	D*****	*****SL**	*N*****N	***G*****LH	*D*****LQK*	FL*****DP*P
	181	SPSSILSTLD	VELPPVCCDI	DI	(SEQ.ID.NO.:4)		
	181	VSKHSVDPG	C		(SEQ.ID.NO.:64)		
	181	*L***L***	*****S**	**	(SEQ.ID.NO.:67)		
	175	*P*A*****	*****Y*A	**	(SEQ.ID.NO.:14)		
	178	*P*A**A**	*****Y*A	**	(SEQ.ID.NO.:2)		

FIG. 24

1012290* 92E45860

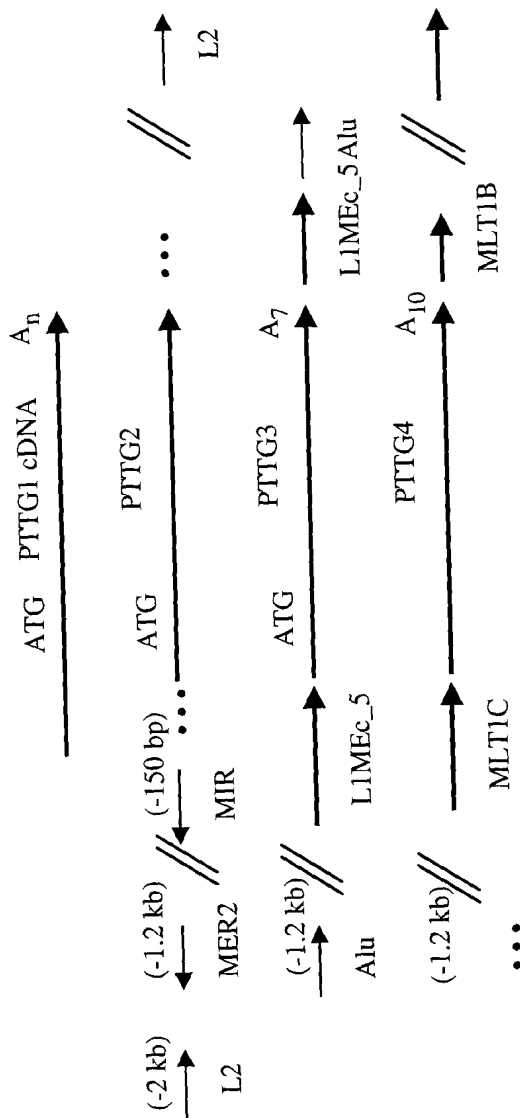


FIG. 25

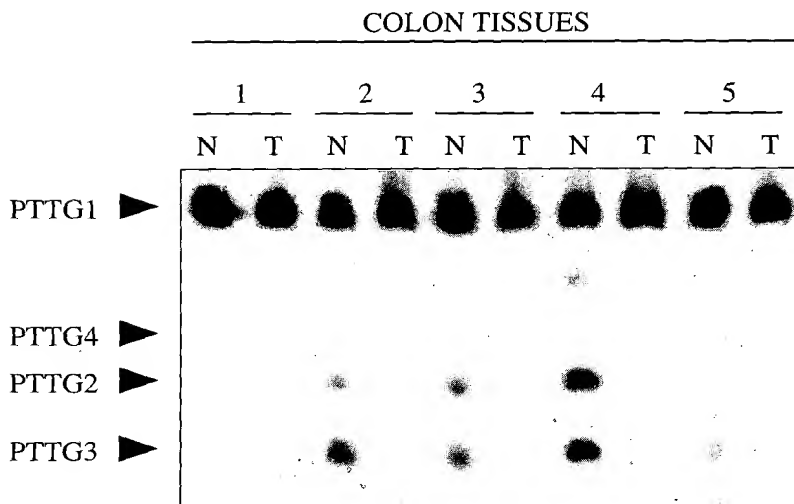


FIG. 27A

0054326-000101
101200-224580

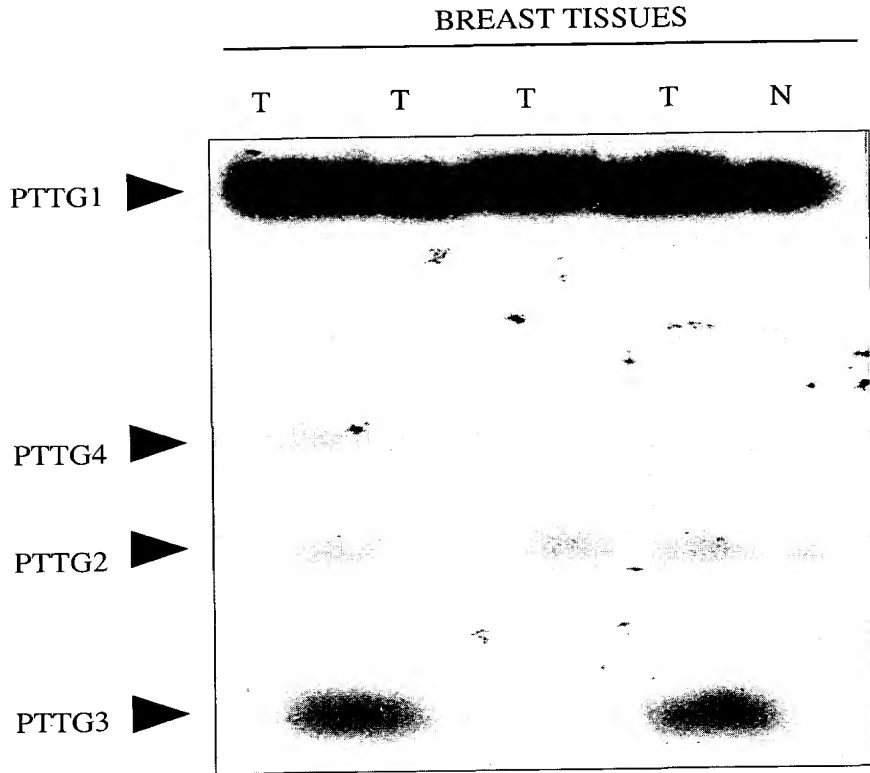


FIG. 27B

107260 92E45860

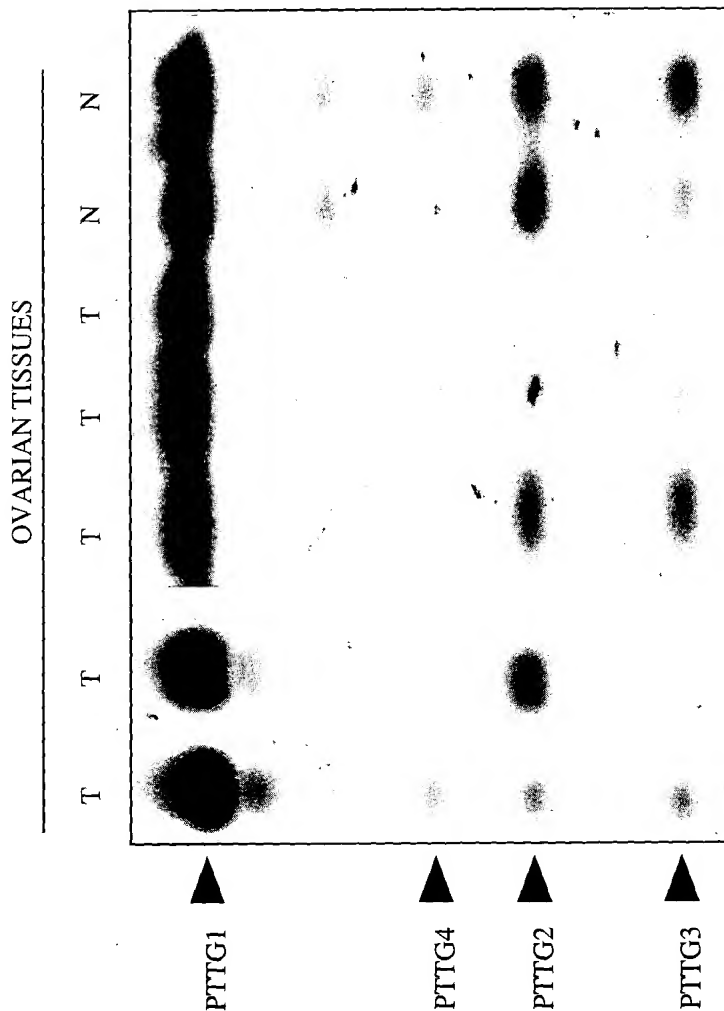


FIG. 27C

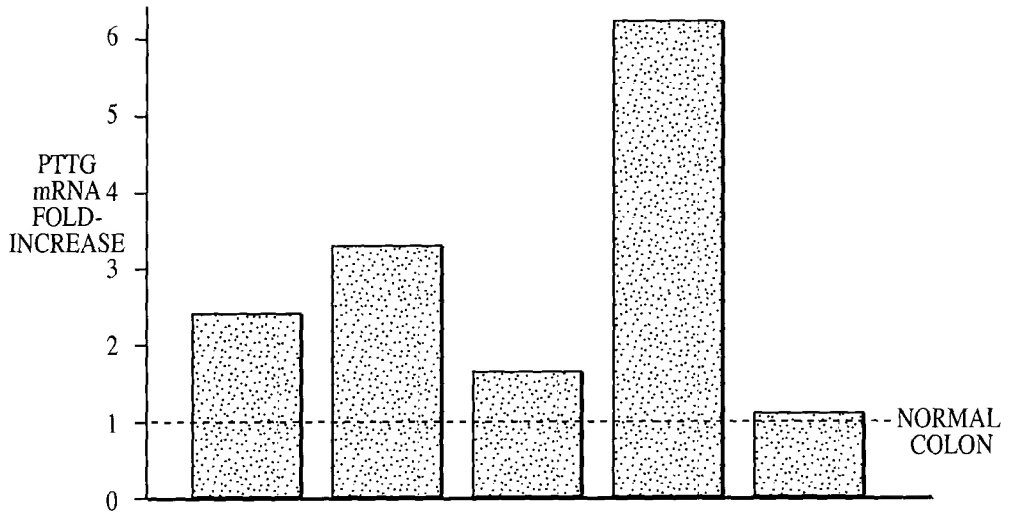


FIG. 28

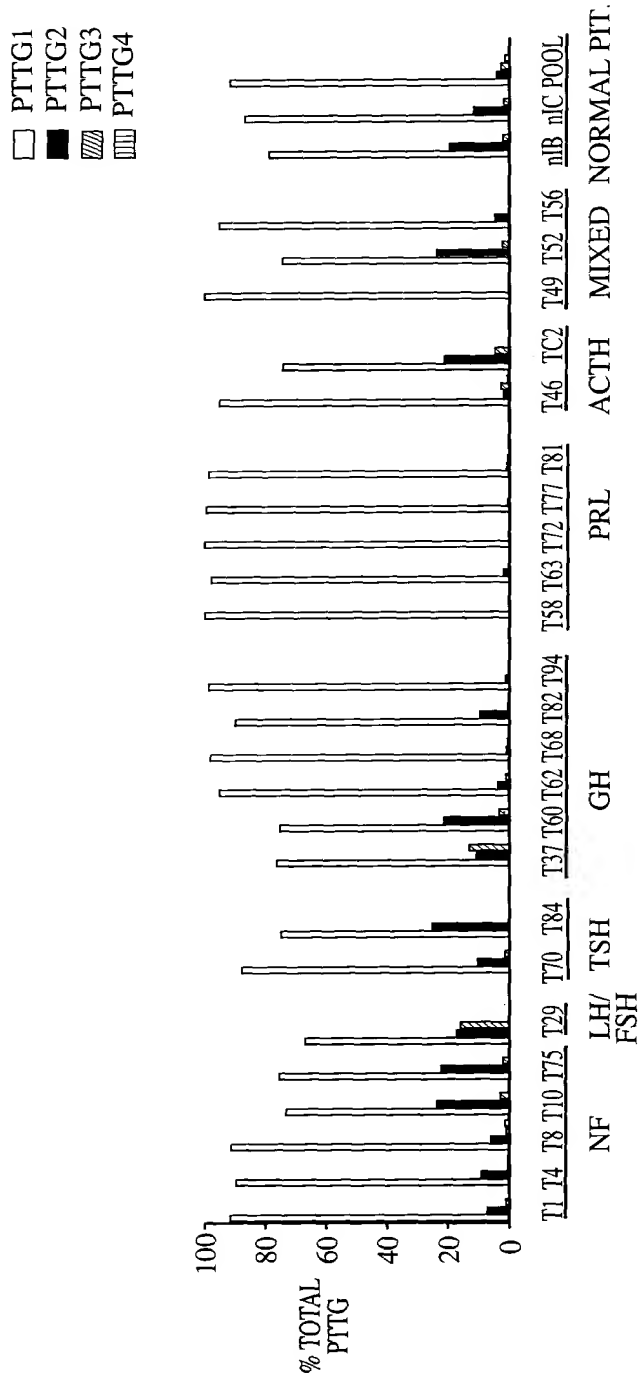


FIG. 29

INTERSPECIFIC HYBRIDS

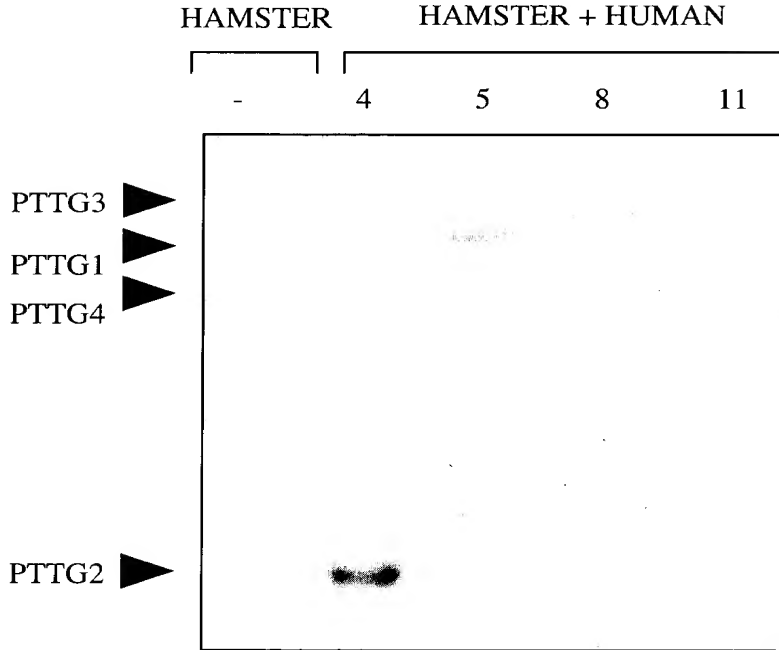


FIG. 30A

COLON TISSUES

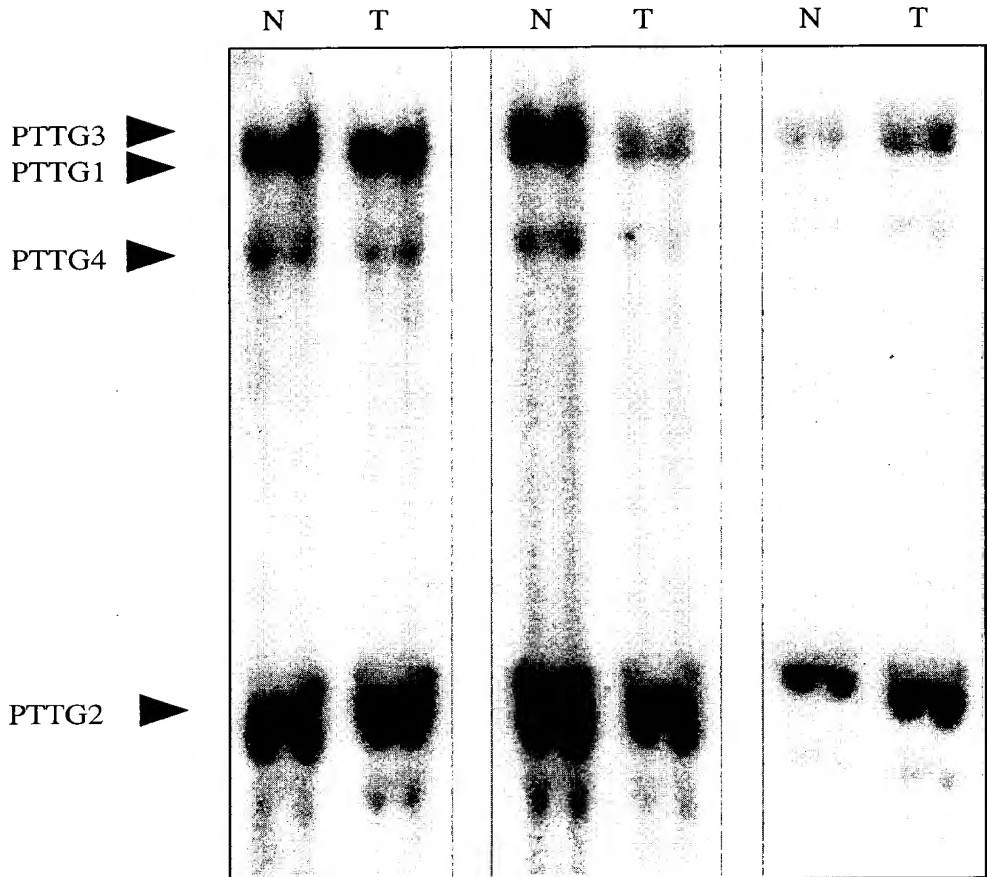


FIG. 30B

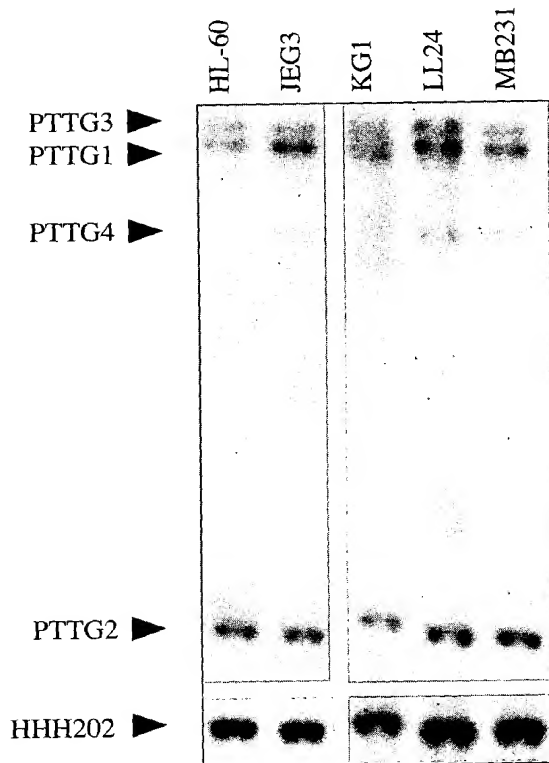


FIG. 30C

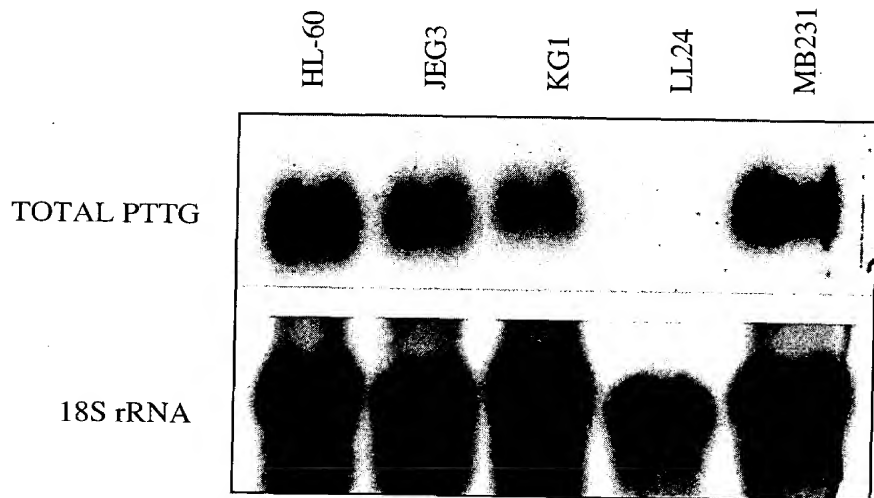


FIG. 30D

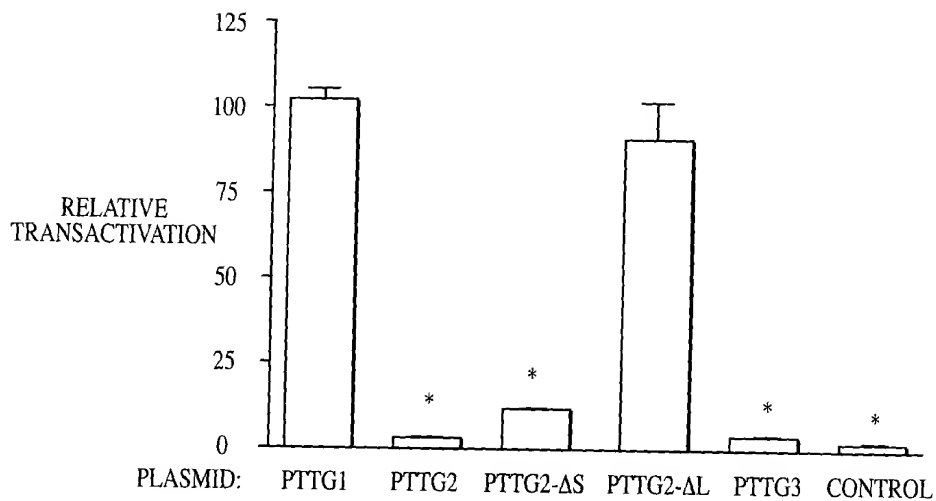


FIG. 31

09854326-022101

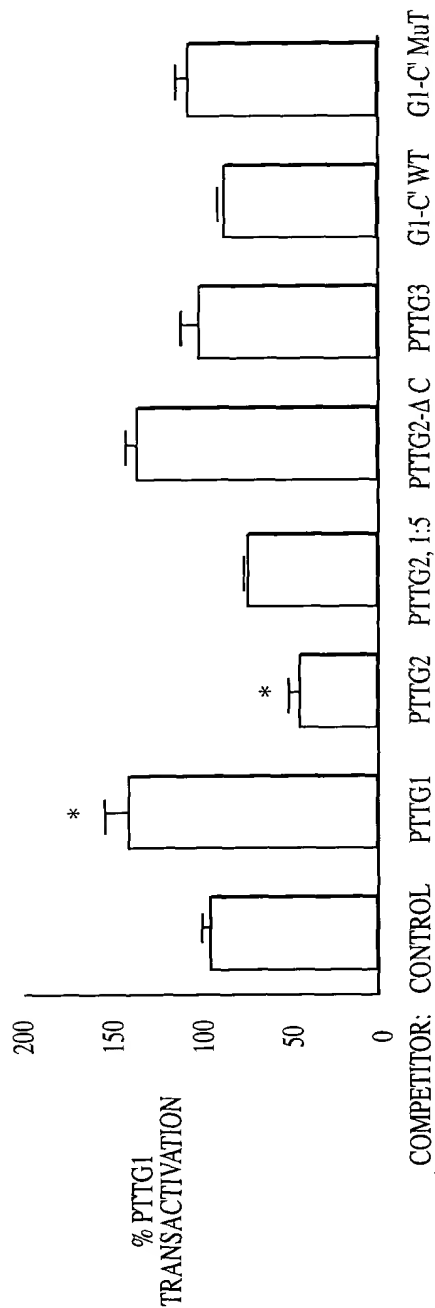


FIG. 32A

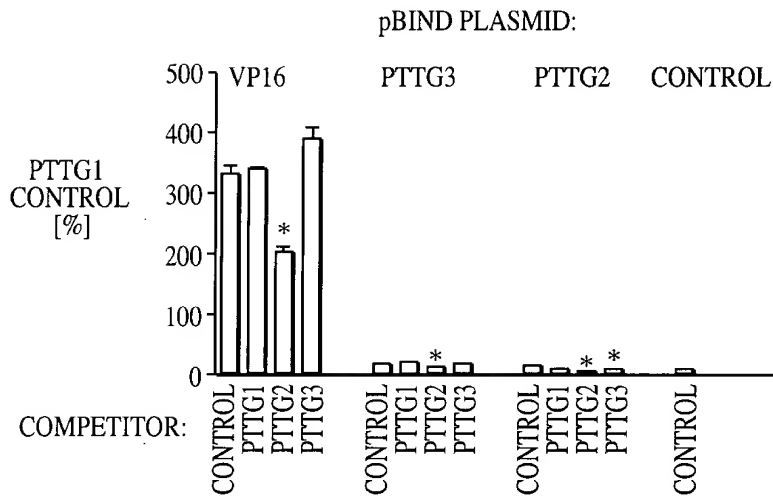


FIG. 32B